



RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number: US/09/762,283 A

Source: PCT

Date Processed by STIC: 11-23-04

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT
MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

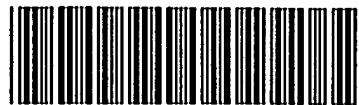
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1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
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Revised 05/17/04

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PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/762,283A

DATE: 11/23/2004

TIME: 12:23:24

Input Set : A:\PTO.YF.txt
 Output Set: N:\CRF4\11232004\I762283A.raw

3 <110> APPLICANT: Perez Gomariz, Rosa
 4 Leceta Martinez, Javier
 5 Delgado Mora, Mario
 6 Martenez Mora, Carmen
 7 <120> TITLE OF INVENTION: Method for Treating Endotoxic Shock and Inflammatory and
 8 Autoimmune Diseases in Mammals
 9 <130> FILE REFERENCE: 100700-09144 [HERR 18.313]
 10 <140> CURRENT APPLICATION NUMBER: US 09/762,283A
 C--> 11 <141> CURRENT FILING DATE: 2004-03-28
 12 <160> NUMBER OF SEQ ID NOS: 3

ERRORED SEQUENCES

38 <210> SEQ ID NO: 3
 39 <211> LENGTH: 27
 40 <212> TYPE: PRT
 41 <213> ORGANISM: Homo sapiens
 42 <400> SEQUENCE: 3
 43 His Ser Asp Gly Ile Phe Thr Asp Ser Tyr Ser Arg Tyr
 44 1 5 10
 E--> 45 Arg Lys Gln Met Ala Val Lys Lys Thy Leu Ala Ala Val
 46 15 20 25
 47 Leu
 E--> 50
 E--> 51
 E--> 52 (continued from previous page . . .)
 E--> 53 (continued)
 E--> 57 11189877.01

Does Not Comply
 Corrected Diskette Needed
 (PG. 1-2)

Invalid Amino Acid Designator

Thy

Delete

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/762,283A

DATE: 11/23/2004
TIME: 12:23:25

Input Set : A:\PTO.YF.txt
Output Set: N:\CRF4\11232004\I762283A.raw

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:45 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1
L:50 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3 ✓
L:50 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1 ✓
M:332 Repeated in SeqNo=3
L:51 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1 ✓
L:52 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:7 ✓✓
L:53 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1
L:57 M:252 E: No. of Seq. differs, <211> LENGTH:Input:27 Found:37 SEQ:3 ✓✓